## **Listing of Claims**

1. (Original) A method of producing a protein with an increased activity or stability, comprising:

replacing an arginine residue capable of being ADP-ribosylated with a tryptophan residue or a phenylalanine residue in a position of an amino acid sequence of the protein, thereby producing the protein with increased activity or stability.

- 2. (Original) The method of claim 1, wherein the protein has an increased antimicrobial activity.
- 3. (Original) The method of claim 2, wherein the antimicrobial activity comprises chemotaxis of T cells, neutrophil recruitment, or cytokine release.
- 4. (Original) The method of claim 3, wherein the cytokine release comprises interleukin-8 release.
  - 5. (Original) The method of claim 2, wherein the protein is a defensin.
  - 6. (Original) The method of claim 5, wherein the defensin is an alpha defensin.
- 7. (Original) The method of claim 2, wherein the arginine residue is substituted in the amino acid sequence of the protein with a tryptophan residue.
- 8. (Original) The method of claim 2, wherein the arginine residue is substituted in the amino acid sequence of the protein with a phenylalanine residue.
- 9. (Original) The method of claim 2, wherein the activity is increased as compared to a polypeptide having an arginine residue in the position of the amino acid sequence of the protein.
- 10. (Original) The method of claim 2, wherein the stability is increased as compared to a polypeptide having an arginine residue in the position of the amino acid sequence of the protein.

- 11. (Original) The method of claim 2, wherein the increased activity or stability is a 100% increase, or a 100% decrease, as compared to a control polypeptide.
- 12. (Original) The method of claim 2, wherein the increased activity or stability is a 50% increase, or a 50% decrease, as compared to a control polypeptide.
- 13. (Currently amended) A method of determining if a protein can be stabilized, comprising:

determining if identifying an arginine residue in the protein is-capable of being ADP-ribosylated; wherein substitution of the arginine residue with a tryptophan or phenylalanine residue increases the stability of the protein, thereby determining if the protein can be stabilized.

- 14. (Original) The method of claim 13, wherein the protein has an antimicrobial activity when administered to a subject.
- 15. (Original) The method of claim 14, wherein determining if an arginine residue in the protein is capable of being ADP-ribosylated comprises:

contacting the protein with an ADP-ribosyltransferase capable of ADP-ribosylating the arginine residue;

measuring an electrophoretic mobility of the protein that was in contact with the ADP-ribosyltransferase; and

comparing the electrophoretic mobility of the protein to an electrophoretic mobility of a first control, wherein a decrease in electrophoretic mobility of the protein, compared to the first control, is an indication that the protein is ADP-ribosylated, thereby determining if the arginine residue in the protein is capable of being ADP-ribosylated.

16. (Original) The method of claim 14, wherein the antimicrobial activity comprises chemotaxis of T cells, neutrophil recruitment or cytokine release.

- 17. (Original) The method of claim14, wherein the protein is a defensin.
- 18. (Original) The method of claim 17, wherein the defensin is an alpha defensin.
- 19. (Original) A composition comprising, a polypeptide comprising an amino acid sequence wherein at least one arginine residue capable of being ADP-ribosylated is substituted with a tryptophan or a phenylalanine residue, wherein the substitution increases the activity or stability of the polypeptide.
- 20. (Original) The composition of claim 21, wherein the polypeptide has an antimicrobial activity.
- 21. (Original) The composition of claim 20, wherein the arginine residue is substituted with a tryptophan residue.
- 22. (Original) The composition of claim 20, wherein the arginine residue is substituted with a phenylalanine residue.
- 23. (Original) The composition of claim 20, wherein the antimicrobial activity comprises chemotaxis of T cells, neutrophil recruitment, or cytokine release.
  - 24. (Original) The composition of claim 20, wherein the protein is a defensin.
  - 25. (Original) The composition of claim 24, wherein the defensin is an alpha defensin.
- 26. (Original) A pharmaceutical composition comprising a therapeutically effective amount of a defensin comprising at least one arginine residue that is substituted by a tryptophan or a phenylalanine residue.
- 27. (Original) The pharmaceutical composition of claim 26, wherein the defensin has antimicrobial activity.

- 28. (Original) The pharmaceutical composition of claim 27, wherein the antimicrobial activity comprises chemotaxis of T cells, neutrophil recruitment or cytokine release.
- 29. (Original) A method of increasing the activity or stability of a defensin polypeptide comprising an arginine residue capable of being ADP-ribosylated, comprising substituting the arginine residue with a tryptophan or a phenylalanine, thereby increasing the activity or the stability of the defensin polypeptide.
- 30. (Original) The method of claim 29, wherein the defensin polypeptide is an alpha defensin.
  - 31. (Original) The method of claim 29, wherein the activity is an antimicrobial activity.
- 32. (Original) The method of claim 31, wherein the antimicrobial activity comprises T cell chemotaxis, neutrophil recruitment, or cytokine release.
- 33. (Original) A method of increasing an immune response in a subject, comprising administering to the subject a therapeutically effective amount of a defensin polypeptide comprising an amino acid substitution, wherein the amino acid substitution is replacement of an arginine capable of being ribosylated with a tryptophan or a phenylalanine, thereby modifying the immune response in the subject.
- 34. (Original) The method of claim 33, wherein the immune response comprises T cell chemotaxis, neutrophil recruitment, or cytokine release.
  - 35. (Original) The method of claim 33, wherein the subject has an immune disorder.